

FORM PTO-1449 <u>INFORMATION DISCLOSURE STATEMENT</u>	ATTY. DOCKET NO.	APPLICATION NO.
	16517.301	10/626,717
	APPLICANTS	
	Scott E. ANDERSEN <i>et al.</i>	
	FILING DATE	GROUP
	July 25, 2003	1634

U.S. PATENT DOCUMENTS

EXAMINER INITIAL	DOCUMENT NUMBER	DATE	NAME	CLASS	SUB- CLASS	FILING DATE
DA1						


FOREIGN PATENT DOCUMENTS

EXAMINER INITIAL	DOCUMENT NUMBER	DATE	COUNTRY	CLASS	SUB- CLASS	TRANSLATION
DB1						Yes No

OTHER (Including Author, Title, Date, Pertinent Pages, etc.)

/JS/	DC1	NCBI Website, BLASTN Accession No. AF020203.1 dated May 15, 1998
↓	DE1	NCBI Website, BLASTN Accession No. U91339.1 dated August 2, 1997
	DE1	NCBI Website, BLASTN Accession No. D38753.1 dated April 29, 2009
	DF1	NCBI Website, BLASTN Accession No. M61109.1 dated April 27, 1993
	DG1	NCBI Website, BLASTN Accession No. L34346.1 dated March 18, 1996
	DH1	NCBI Website, BLASTX Accession No. AAB65144 dated August 2, 1997
	DI1	NCBI Website, BLASTX Accession No. P22243.1 dated June 16, 2009
	DJ1	NCBI Website, BLASTX Accession No. BAA08636 dated April 28, 2009
↓	DK1	NCBI Website, BLASTX Accession No. AAB65145.1 dated August 2, 1997
	DL1	NCBI Website, BLASTX Accession No. Q96456.1 dated June 16, 2009

EXAMINER	/Jehanne Sitton/	DATE CONSIDERED	12/30/2009
EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to Applicant.			

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	DM1	Tian <i>et al.</i> , "How Well is Enzyme Function Conserved as a Function of Pairwise Sequence Identity?", <i>J. Mol. Biol.</i> 333:863-882 (October 31, 2003)	
	DN1	Todd <i>et al.</i> , "Evolution of protein function, from a structural perspective", <i>Curr Opin Chem Biol.</i> 3(5):548-56 (October 1999) (abstract only)	
	DO1	Skolnick <i>et al.</i> , "From genes to protein structure and function: novel applications of computational approaches in the genomic era", <i>Trends Biotechnol.</i> 18(1):34-9 (January 2000)	
	DP1	Wilson <i>et al.</i> , "Assessing annotation transfer for genomics: quantifying the relations between protein sequence, structure and function through traditional and probabilistic scores", <i>J. Mol. Biol.</i> 297(1):233-249 (March 17, 2000) (abstract only)	

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